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AMENDMENTS TO THE CLAIMS

Claims 1-4. Canceled

[c5]-5. (Currently Amended) The isolated nucleic acid of Claim 1—An isolated nucleic acid having at least 99% nucleic acid sequence identity to:

- (a) a nucleic acid sequence encoding the polypeptide shown in Figure 6 (SEQ ID NO:6);
- (b)—a nucleic acid-sequence encoding the polypeptide shown in Figure 6 (SEQ ID NO:6), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 6 (SEQ ID NO:6);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 6 (SEQ ID NO:6), lacking its associated signal peptide;
- (e) the nucleic acid sequence of shown in Figure 5 (SEQ ID NO:5); wherein said nucleic acid encodes a polypeptide which stimulates TNF-α release from human blood.
- (f) the full-length coding sequence of the nucleic acid sequence shown in Figure 5 (SEQ ID NO:5); or
- (g)—the full-length coding sequence of the cDNA deposited under ATCC accession number 209399.

[c6] 6. (Currently Amended) An isolated nucleic acid comprising:

- (a) a nucleic acid sequence encoding the polypeptide of shown in Figure 6 (SEQ ID NO:6);
- (b) a nucleic acid sequence encoding the polypeptide of shown in Figure 6 (SEQ ID NO:6), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 6 (SEQ ID NO:6);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 6 (SEQ ID NO:6), lacking its associated signal peptide;
 - (e) the nucleic acid sequence of shown in Figure 5 (SEQ ID NO:5);
- (f) the full-length coding sequence of the nucleic acid sequence of shown in Figure 5-(SEQ ID NO:5); or

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(g) the full-length coding sequence of the cDNA deposited under ATCC accession number 209399;

wherein said extracellular domain is amino acids 17-234 of SEQ ID NO:6.

- [e7]-7. (Currently Amended) The isolated nucleic acid of Claim 6 comprising a nucleic acid sequence encoding the polypeptide of shown in Figure 6 (SEQ ID NO:6).
- [c8] 8. (Currently Amended) The isolated nucleic acid of Claim 6 comprising a nucleic acid sequence encoding the polypeptide of shown in Figure 6 (SEQ ID NO:6), lacking its associated signal peptide.
- [c9] 9. (Currently Amended) The isolated nucleic acid of Claim 6 comprising a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 6 (SEQ ID NO:6), wherein said extracellular domain is amino acids 17-234 of SEQ ID NO:6.
- [c10]-10. (Currently Amended) The isolated nucleic acid of Claim 6 comprising a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 6 (SEQ ID NO:6), lacking its associated signal peptide, wherein said extracellular domain is amino acids 17-234 of SEQ ID NO:6.
- [c11] 11. (Currently Amended) The isolated nucleic acid of Claim 6 comprising the nucleic acid sequence of shown in Figure 5 (SEQ ID NO:5).
- [c12]-12. (Currently Amended) The isolated nucleic acid of Claim 6 comprising the full-length coding sequence of the nucleic acid sequence of shown in Figure 5 (SEQ ID NO:5).
- [c13]-13. (Currently Amended) The isolated nucleic acid of Claim 6 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 209399.

Claims 14-16. Canceled

- [c17] 17. (Currently Amended) A vector comprising the nucleic acid of Claim-16.
- [c18]-18. (Currently Amended) The vector of Claim 17, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.

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[c19] 19. (Currently Amended) A—An isolated host cell comprising the

vector of Claim 17.

[c20]-20. (Currently Amended) The host cell of Claim 19, wherein said cell

is a CHO cell, an E. coli or a yeast cell.

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DELETION OF INVENTORS

Please correct the inventorship under 37 CFR §1.48(b) by removing the following inventors from the present application:

Dan L. Eaton, Ellen Filvaroff, Mary E. Gerritsen, Christopher J. Grimaldi and Colin K. Watanabe.